

GenCore version 5.1.6
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OM protein - protein search, using SW model

Run on: May 10, 2004, 07:26:27 ; Search time 23 Seconds
(without alignments)
1335.541 Million cell updates/sec

Title: US-09-943-446a-6

Perfect score: 3169

Sequence: 1 MEAVRRLAPLALLCPVL.....EASAPRRPALLQEMETW 595

Scoring table: BLOSUM62

Gapop 10.0 , Gapext 0.5

Searched: 389414 seqs, 51625971 residues

Total number of hits satisfying chosen parameters: 389414

Minimum DB seq length: 0
Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0%

Listing first 45 summaries

Database :
1: Issued Patents AA:*
2: /cgn2_6/prodata/2/iaa/5A_COMB.pep:*
3: /cgn2_6/prodata/2/iaa/5B_COMB.pep:*
4: /cgn2_6/prodata/2/iaa/5A_COMB.pep:*
5: /cgn2_6/prodata/2/iaa/5B_COMB.pep:*
6: /cgn2_6/prodata/2/iaa/backfill1.pep:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

| Result No. | Score | Query Match | Length DB | ID | Description |
|------------|--------|-------------|-----------|----|--------------------|
| 1 | 3020 | 95.3 | 593 | 2 | US-08-468-249A-21 |
| 2 | 2819 | 89.0 | 591 | 2 | US-08-468-249A-20 |
| 3 | 2444 | 77.1 | 585 | 2 | US-08-142-551B-125 |
| 4 | 2444 | 77.1 | 585 | 2 | US-08-468-249A-19 |
| 5 | 2440 | 77.0 | 585 | 1 | US-08-142-439A-6 |
| 6 | 2440 | 77.0 | 585 | 2 | US-08-468-249A-18 |
| 7 | 2286.5 | 72.2 | 515 | 2 | US-08-468-249A-18 |
| 8 | 1923.5 | 60.7 | 536 | 4 | US-09-449-632-2 |
| 9 | 1513.5 | 47.8 | 532 | 4 | US-09-449-632-4 |
| 10 | 1385 | 43.7 | 575 | 4 | US-09-449-632-5 |
| 11 | 1345.5 | 42.5 | 541 | 3 | US-08-468-011A-2 |
| 12 | 1345.5 | 42.5 | 541 | 4 | US-09-236-468A-2 |
| 13 | 1345.5 | 42.5 | 541 | 5 | PCT-US95-07085-2 |
| 14 | 994 | 31.4 | 207 | 4 | US-08-811-519-31 |
| 15 | 863 | 27.2 | 449 | 1 | US-08-142-439A-5 |
| 16 | 863 | 27.2 | 449 | 2 | US-08-869-477-5 |
| 17 | 855 | 27.0 | 1324 | 2 | US-08-811-897A-56 |
| 18 | 855 | 27.0 | 1324 | 4 | US-09-201-474-17 |
| 19 | 824 | 26.0 | 458 | 1 | US-08-112-817C-2 |
| 20 | 782.5 | 24.7 | 437 | 2 | US-08-538-816A-2 |
| 21 | 782.5 | 24.7 | 437 | 2 | US-09-076-651-2 |
| 22 | 782.5 | 24.7 | 437 | 4 | US-09-208-394-2 |
| 23 | 765.5 | 24.2 | 431 | 2 | US-08-538-816A-9 |
| 24 | 765.5 | 24.2 | 431 | 2 | US-09-076-651-9 |
| 25 | 765.5 | 24.2 | 431 | 4 | US-09-208-394-9 |
| 26 | 765.5 | 24.2 | 438 | 2 | US-08-538-816A-1 |
| 27 | 765.5 | 24.2 | 438 | 2 | US-09-076-651-1 |

| | | | | | | |
|----|-------|------|-----|---|-------------------|-------------------|
| 28 | 765.5 | 24.2 | 438 | 4 | US-09-208-394-1 | Sequence 1, Appl |
| 29 | 764.5 | 24.1 | 525 | 2 | US-08-811-897A-23 | Sequence 23, Appl |
| 30 | 764.5 | 24.1 | 525 | 2 | US-08-855-213-23 | Sequence 23, Appl |
| 31 | 764.5 | 24.1 | 525 | 4 | US-09-201-474-23 | Sequence 23, Appl |
| 32 | 754 | 23.8 | 448 | 2 | US-08-811-897A-22 | Sequence 22, Appl |
| 33 | 754 | 23.8 | 448 | 2 | US-08-855-213-22 | Sequence 22, Appl |
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| 35 | 753.5 | 23.7 | 485 | 4 | US-08-811-897A-17 | Sequence 17, Appl |
| 36 | 752.5 | 23.7 | 485 | 2 | US-08-855-213-17 | Sequence 17, Appl |
| 37 | 752.5 | 23.7 | 485 | 4 | US-09-201-474-17 | Sequence 17, Appl |
| 38 | 752 | 23.7 | 552 | 2 | US-08-811-897A-27 | Sequence 27, Appl |
| 39 | 752 | 23.7 | 552 | 2 | US-08-855-213-27 | Sequence 27, Appl |
| 40 | 752 | 23.7 | 552 | 4 | US-09-201-474-27 | Sequence 27, Appl |
| 41 | 751.5 | 23.7 | 553 | 2 | US-08-811-897A-25 | Sequence 25, Appl |
| 42 | 751.5 | 23.7 | 553 | 2 | US-08-855-213-25 | Sequence 25, Appl |
| 43 | 751.5 | 23.7 | 553 | 4 | US-09-201-474-25 | Sequence 25, Appl |
| 44 | 750.5 | 23.7 | 467 | 2 | US-08-811-897A-19 | Sequence 19, Appl |
| 45 | 750.5 | 23.7 | 467 | 2 | US-08-855-213-19 | Sequence 19, Appl |

ALIGNMENTS

RESULT 1
US-08-468-249A-21
Sequence 21, Application US/08468249A
Patent No. 5886148

GENERAL INFORMATION:

APPLICANT: Segre et al., Gino V.
TITLE OF INVENTION: PARATHYROID HORMONE RECEPTOR AND DNA
TITLE OF INVENTION: ENCODING SAME
NUMBER OF SEQUENCES: 21
CORRESPONDENCE ADDRESS:
ADDRESS: Fish & Richardson P.C.
CITY: Boston
STATE: MA
COUNTRY: USA
ZIP: 02110-2804
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/468,249A
FILING DATE: 06-JUN-1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/864,475
FILING DATE: 06-APR-1992

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/681,702
FILING DATE: 04-MAY-1991
ATTORNEY/AGENT INFORMATION:
NAME: Fraser, Janis K.

REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 00786/071003
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 593 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein

US-08-468-249A-21
Query Match 95.3%; Score 3020; DB 2; Length 593;
Best Local Similarity 95.2%; Pred. No. 4.6e-278;
Matches 570; Conservative 5; Mismatches 14; Indels 10; Gaps 3;

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QY 1 MGAARIAPGALILCCPYLSAVALVDADVMTKEBQIFILHRAOAOCCOKLKEVLOPA 60
DB 1 MGAARIAPGALILCCPYLSAVALVDADVMTKEBQIFILHRAOAOCCOKLKEVLOPA 60
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DB 61 DIMESDKGMASTSGKPKKEKASGKLYPSEEDKEVPTGSRHGRPCLEPMDHILCWL 120
QY 61 SIMESEKGMATSGKPKKEKASGKLYPSEEDKEVPTGSRHGRPCLEPMDHILCWL 120
DB 61 SIMESEKGMATSGKPKKEKASGKLYPSEEDKEVPTGSRHGRPCLEPMDHILCWL 120
QY 121 GAGEVAVAPCPDITVDNHNHGHAYRRCDSNGSMELVPGHNRWTANYSCEVKTNETRE 180
DB 121 GAGEVAVAPCPDITVDNHNHGHAYRRCDSNGSMELVPGHNRWTANYSCEVKTNETRE 180
QY 121 GAGEVAVAPCPDITVDNHNHGHAYRRCDSNGSMELVPGHNRWTANYSCEVKTNETRE 180
DB 121 GAGEVAVAPCPDITVDNHNHGHAYRRCDSNGSMELVPGHNRWTANYSCEVKTNETRE 180
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DB 181 REVDRLGMITVGVSYSLATVAVLILAFRLHCTRYNIHMLFLSFLRAVSIFFVK 240
QY 241 DAVISGATLDEARLTREELRAIAQAPPPATPAAGCRAVAVTFEFLATNTYWI 299
DB 241 DAVISGATLDEARLTREELRAIAQAPPPATPAAGCRAVAVTFEFLATNTYWI 299
QY 241 DAVISGATLDEARLTREELRAIAQAPPPATPAAGCRAVAVTFEFLATNTYWI 300
DB 241 DAVISGATLDEARLTREELRAIAQAPPPATPAAGCRAVAVTFEFLATNTYWI 300
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QY 301 VEGYLHSLIFMAFSEKXYLMGFTVGMGLPAVFVAWVSVAATLANTGCMDSGK 360
DB 301 VEGYLHSLIFMAFSEKXYLMGFTVGMGLPAVFVAWVSVAATLANTGCMDSGK 360
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DB 420 YIVFMATPYEVSGTLMOVQNHMELFNSFOGFVAIIFCGNGEVOAEIKKMSRWTLA 480
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DB 481 LDFRKARSSSSSYSGPMVSHTSVTNVPRAAGLPLSPRLPAAATTTATTNGHPPI 534
QY 540 PGHTKPGAP---TLPATPPATAAPKXDDGFLNCSGGLDEASAPRPPALLOEWEVTV 595
DB 540 PGHTKPGAP---TLPATPPATAAPKXDDGFLNCSGGLDEASAPRPPALLOEWEVTV 593
QY 595 PGHTKPGAP---TLPATPPATAAPKXDDGFLNCSGGLDEASAPRPPALLOEWEVTV 595
DB 595 PGHTKPGAP---TLPATPPATAAPKXDDGFLNCSGGLDEASAPRPPALLOEWEVTV 593

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NAME: Fraser, Janis K.
REGISTRATION NUMBER: 34,819
REFERENCE/DOCKET NUMBER: 00786/071003
TELEPHONE: 617/542-5070
TELEFAX: 617/542-8906
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 591 amino acids
TYPE: amino acid
TOPOLOGY: linear
MOLECULE TYPE: protein
US-08-468-249A-20

Query Match
Best Local Similarity 89.0%; Score 2819; DB 2; Length 591;
Matches 530; Conservative 21; Mismatches 34; Indels 16; Gaps 4;

1 MGAARIAPGALILCCPYLSAVALVDADVMTKEBQIFILHRAOAOCCOKLKEVLOPA 60
1 MGAARIAPGALILCCPYLSAVALVDADVMTKEBQIFILHRAOAOCCOKLKEVLOPA 60
61 DIMESDKGMASTSGKPKKEKASGKLYPSEEDKEVPTGSRHGRPCLEPMDHILCWL 120
61 DIMESDKGMASTSGKPKKEKASGKLYPSEEDKEVPTGSRHGRPCLEPMDHILCWL 120
61 NIMESDKGMASTSGKPKKEKASGKLYPSEEDKEVPTGSRHGRPCLEPMDHILCWL 120
121 GAGEVAVAPCPDITVDNHNHGHAYRRCDSNGSMELVPGHNRWTANYSCEVKTNETRE 180
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121 GAGEVAVAPCPDITVDNHNHGHAYRRCDSNGSMELVPGHNRWTANYSCEVKTNETRE 180
181 REVDRLGMITVGVSYSLATVAVLILAFRLHCTRYNIHMLFLSFLRAVSIFFVK 240
181 REVDRLGMITVGVSYSLATVAVLILAFRLHCTRYNIHMLFLSFLRAVSIFFVK 240
241 DAVISGATLDEARLTREELRAIAQAPPPATPAAGCRAVAVTFEFLATNTYWI 299
241 DAVISGATLDEARLTREELRAIAQAPPPATPAAGCRAVAVTFEFLATNTYWI 299
241 DAVISGATLDEARLTREELRAIAQAPPPATPAAGCRAVAVTFEFLATNTYWI 300
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360 WIIQVPIIASIVNFIINIVRLATKRETNAGRCTROQYRKLIKSTVIMPLFGVH 419
361 WIIQVPIIASIVNFIINIVRLATKRETNAGRCTROQYRKLIKSTVIMPLFGVH 420
420 YIVFMATPYEVSGTLMOVQNHMELFNSFOGFVAIIFCGNGEVOAEIKKMSRWTLA 479
421 YIVFMATPYEVSGTLMOVQNHMELFNSFOGFVAIIFCGNGEVOAEIKKMSRWTLA 480
480 YIVFMATPYEVSGTLMOVQNHMELFNSFOGFVAIIFCGNGEVOAEIKKMSRWTLA 480
481 LDFRKARSSSSSYSGPMVSHTSVTNVPRAAGLPLSPRLPAAATTTATTNGHPPI 539
481 LDFRKARSSSSSYSGPMVSHTSVTNVPRAAGLPLSPRLPAAATTTATTNGHPPI 533
540 PGHTKPGAP---TLPATPPATAAPKXDDGFLNCSGGLDEASAPRPPALLOEWEVTV 594
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591 M 591
591 M 591

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RESULT 3
US-08-142-551B-125
Sequence 125, Application US/08142551B
Patent No. 5814603
GENERAL INFORMATION:
APPLICANT: Oldenburg, Kevin R.
TITLE OF INVENTION: COMPOUNDS WITH PTH ACTIVITY AND
RECOMBINANT DNA VECTORS ENCODING SAME
NUMBER OF SEQUENCES: 132

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405 May